

Fig. 1

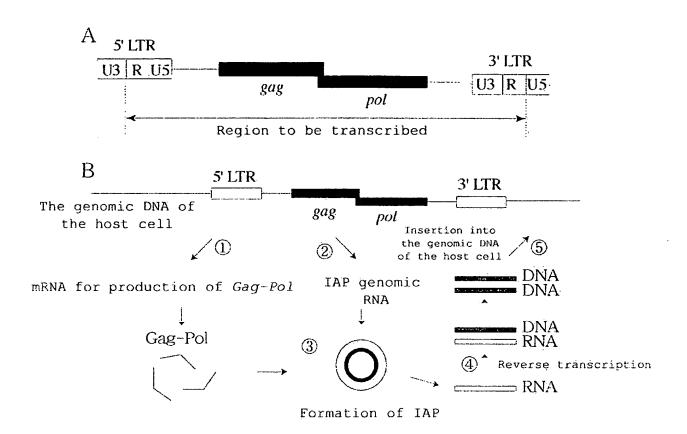
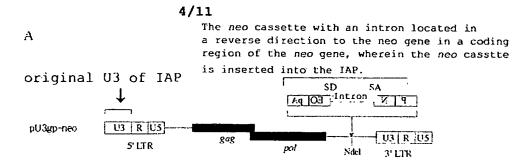
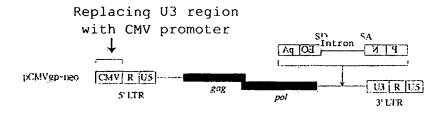


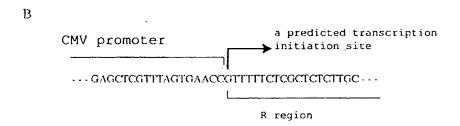
Fig. 2

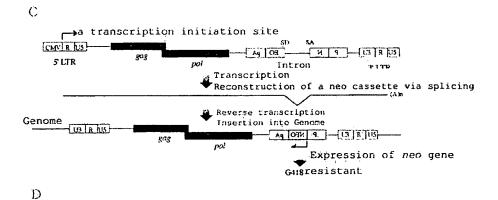
## 3/11 The neo cassette with an intron located in Α a reverse direction to the neo gene in a coding region of the neo gene, wherein the neo (1)Replacing U3 region casstte is inserted into the IAP. with CMV promoter SD SA Intron CMV R U5 gag 5'LTR pol 3' LTR Transcription В ightarrowinitiation site CMV R U5 U3 R U5 gag 5' LTR pol 3'LTR Transcription Reconstruction of a neo cassette via splicing Reverse transcription Insertion into Genome Genome UB R US NEO pA 9 -- U3 R U5 pol Expression of neo gene G418 resistant (i) IAP (ii) IAP-delta pol (iii) LINEI

Fig. 3









Vector The number of G418 resistant colonies

Mock 0 pU3gp-neo 0 pCMVgp-neo 110 pJM101/L1.3(LINE1) 22

Fig. 4

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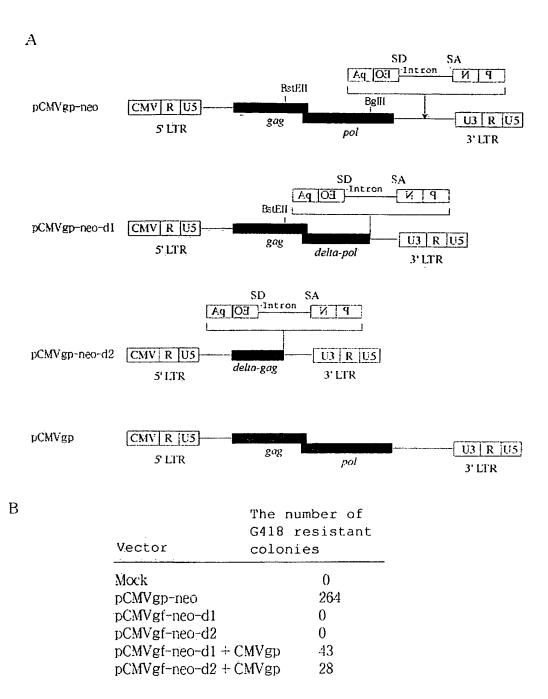


Fig. 5

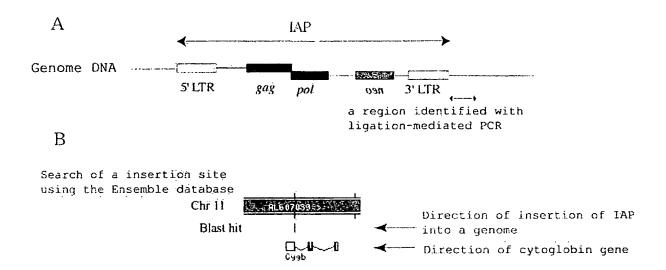
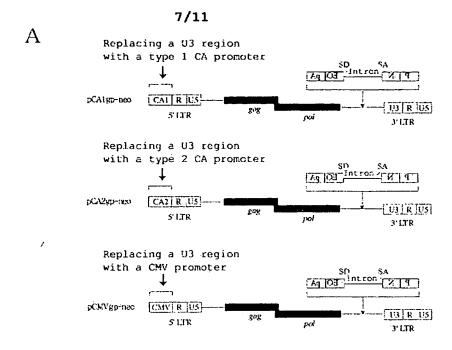
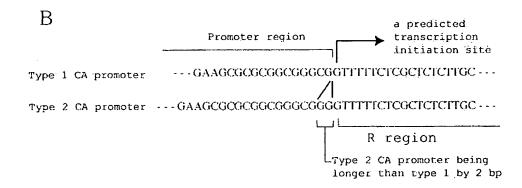


Fig. 6



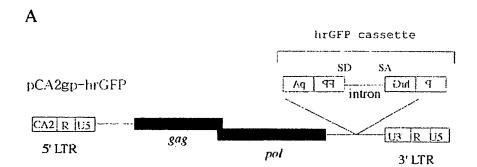


		The number of G418 resistant colonies	
Vector	NIH3T3	Hela	
pCA1gp-neo	65 92	95 223	
pCA2gp-neo	17 <u>3</u> 124	230 185	
pCMVgp=neo	92 90	208 177	

Fig. 7

C

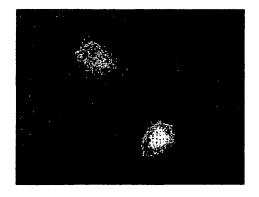
8/11



В



Bright-field



Fluorescence-field

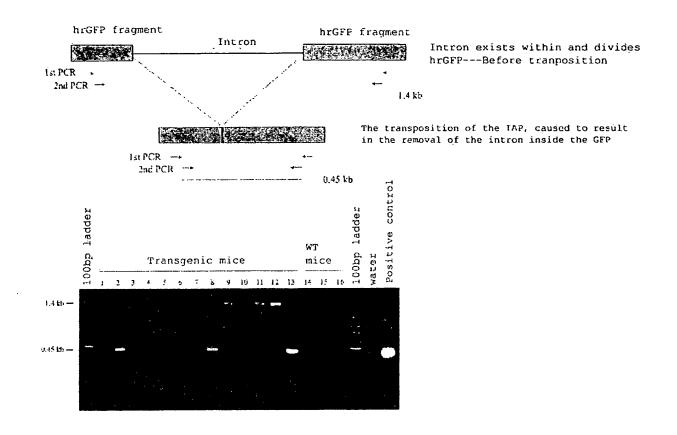
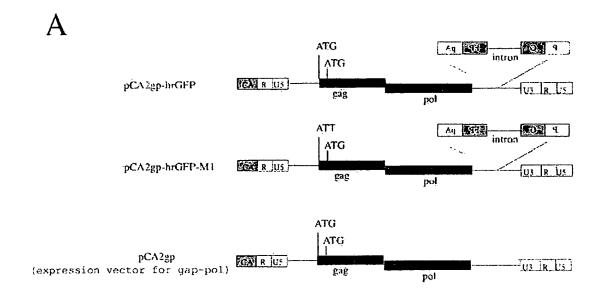


Fig. 9

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The first ATG in a gag gene, which is modified in pCA2gp-hrGFP-M1 into ATT.

The second ATG in a gag gene, from which translation in pCA2gp-hrGFP-M1 is predicted to start.



WT ATG AAT TCA GAA CTT TTC ACC TGG GGA ACG AGA GTA CCA GTG AGT AFG TTT GGC CTT GAA (pCA2gp-hrGFP) Met Asn Ser Glu Leu Phe Ser Trp Gly Thr Arg Val Pro Vul Ser Met Phe Gly Leu Glu

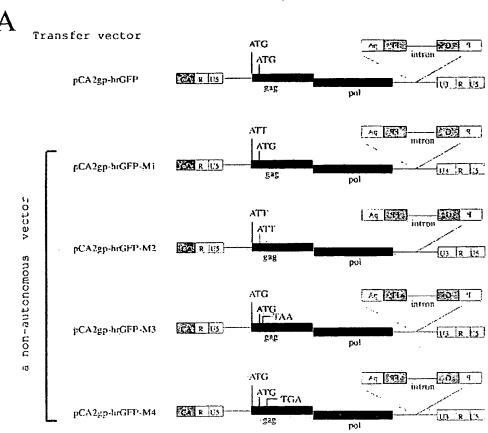
B

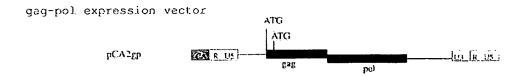
Vector	% GFP-positive	
pCA2gp-hrGFP	0.29	
pCA2gp-hiGFP-M1	0	
pCA2gp-hrGFP-M1 + pCA2gp	0.22	

Fig. 10

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B

Transfer vector	% GFP-positive		
	+pBluescript	+pCA2gp	
pCA2gp-hrGFP	0.29		not done
pCA2gp-hrGFP-M1	0		0.215
pCA2gp-hrGFP-M2	0		0.005
pCA2gp-hrGFP-M3	0		0.04
pCA2gp-hrGFP-M4	0		0.015

Fig. 11